

MILLNVLRICI IVCLVNDGAG KHSEGRERTK TYSLNSRGYF 40
RKERGARRSK ILLVNTKGKD EPHIGHGDFG LVAELFDSTR 80
THTNRKEPDM NKVKLFSTVA HGNKSARRKA YNGSRRNIFS 120
RRSFDFKRNT E VTEKPGAKMF WNNFLVKMNG APQNTSHGSK 160
AQEIMKEACK TLPFTQNIVH ENCDRMVIQN NLCFGKCISL 200
HVPNQQDRRN TCSHCLPSKF TLNHLTNCT GSKNVVKVVM 240
MVEECTCEAH KSNFHQTAQF NMDTSTTLHH 270

Figure 1. Deduced amino acid sequence of Xenopus cerberus protein. SEQ ID NO:1.

Figure 2. Nucleotide sequence of the full-length cerberus DNA derived from the *Xenopus* organizer. The sense strand is on top (in the 5' to 3' direction) and the antisense strand on the bottom line (on the opposite direction). SEQ ID NO:2.

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|---|------|
| GAATTCCAG CAAGTCGCTC AGAACACTG CAGGGTCTAG ATATCATACA ATGTTACTAA CTTAAGGGTC GTTCAGCGAG TCTTGTGAC GTCCCAGATC TATAGTATGT TACAATGATT | 60 |
| ATGTAACTCAG GATCTGTATT ATCGTCTGCC TTGTGAATGA TGGAGCAGGA AAACACTCAG TACATGAGTC CTAGACATAA TAGCAGACGG AACACTTACT ACCTCGCTC TTTGTGAGTC | 120 |
| AAGGACGAGA AAGGACAAAA ACATATTCAC TAAACAGCAG AGGTTACTTC AGAAAAGAAA TCCCTGCTCT TTCCCTGTTT TGTATAAGTG AATTGTCGTC TCCAATGAAG TCTTTCTTT | 180 |
| GAGGAGCACG TAGGAGCAAG ATTCTGCTGG TGAATACTAA AGGTCTTGAT GAACCCCCACA CTCCTCGTGC ATCCTCGTTC TAAGACGACC ACTTATGATT TCCAGAACTA CTTGGGGTGT | 240 |
| TTGGGCATGG TGATTTTCGC TTAGTAGCTG AACTATTTGA TTCCACCAGA ACACATACAA AACCCGTACC ACTAAAAGCG AATCATCGAC TTGATAAACT AAGGTGGTCT TGTGTATGTT | 300 |
| ACAGAAAAGA GCCAGACATG AACAAAGTCA AGCTTTCTC AACAGTTGCC CATGGAAACA TGTCTTTCT CGGTCTGTAC TTGTTTCAGT TCGAAAAGAG TTGTCAACGG GTACCTTTGT | 360 |
| AAAGTGCAAG AAGAAAAGCT TACAATGGTT CTAGAAGGAA TATTTTCCT CGCCGTTCTT TTTCACGTTTC TTCTTTTCGA ATGTTACCAA GATCTTCCTT ATAAAAGGA GCGGCAAGAA | 420 |
| TTGATAAAAG AAATACAGAG GTTACTGAAA AGCCTGGTGC CAAGATGTT TC GGAAACAATT AACTATTTTC TTTATGTCTC CAATGACTTT TCGGACCACG GTTCTACAAG ACCTTGTAA | 480 |
| TTTTGGTTAA AATGAATGGA GCCCCACAGA ATACAAGCCA TGGCAGTAAA GCACAGGAAA AAAACCAATT TTACTTACCT CGGGGTGTCT TATGTTCGGT ACCGTCAATT CGTGTCTTT | 540 |
| TAATGAAAGA AGCTTGCAAA ACCTTGTTTT TCACTCAGAA TATTGTACAT GAAAAGTGTG ATTACTTTCT TCGAACGTTT TGGAACAAAA AGTGAAGTCTT ATAACATGTA CTTTGACAC | 600 |
| ACAGGATGGT GATACTGAAAC AATCTGTGCT TTGGTAAATG CATCTCTC CATGTTCCAA TGTCTTACCA CTATGTCTTG TTAGACACGA AACCAATTAC GTAGAGAGAG GTACAAGGTT | 660 |
| ATCAGCAAGA TCGACGAAAT ACTTGTCCC ATTGCTTGCC GTCCAAATT ACCCTGAACC TAGTCGTTCT AGCTGCTTTA TGAACAAGGG TAACGAACGG CAGGTTAAA TGGGACTTGG | 720 |
| ACCTGACGCT GAATTGTACT GGATCTAAGA ATGTTAGTAA GGTTGTCTG ATGGTAGAGG TGGACTGCGA CTTAACATGA CCTAGATTCT TACATCATT CCAACAGTAC TACCATCTCC | 780 |
| AATGCACGTG TGAAGCTCAT AAGAGCAACT TCCACCAAC TGCACTGTTT AACATGGATA TTACGTGCAC ACTTCGAGTA TTCTCGTTGA AGGTGGTTG ACGTGTCAAA TTGTACCTAT | 840 |
| CATCTACTAC CCTGCACCAT TAAAGGACTG CCATACAGTA TGGAAATGCC CTTTTGTTGG GTAGATGATG GGACGTGGTA ATTTCTGAC GGATGTCTAC ACCTTTACGG GAAAACAACC | 900 |
| AATATTGTT ACATACTATG CATCTAAAGC ATTATGTTGC CTTCTATTTC ATATAACCAC TTATAAACAA TGTATGATAC GTAGATTCG TAATACAACG GAAGATAAAG TATATTGGTG | 960 |
| ATGGAATAAG GATTGTATGA ATTATAATTA ACAAAATGGCA TTTTGTGTA CATGCAAGAT TACCTTATTC CTAACATACT TAATATTAAT TGTTTACCGT AAAACACATT GTACGTTCTA | 1020 |

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|------------|------------|-------------|------------|------------|-------------|------|
| CTCTGTTCCA | TCAGTTGCAA | GATAAAAGGC | AATATTGTT | TGACTTTTT | TCTACAAAAT | 1080 |
| GAGACAAGGT | AGTCAACGTT | CTATTTCCG | TTATAAACAA | ACTGAAAAAA | AGATGTTTA | |
| GAATACCCAA | ATATATGATA | AGATAATGGG | GTCAAAAATG | TTAAGGGGTA | ATGTAATAAT | 1140 |
| CTTATGGGTT | TATATACTAT | TCTATTACCC | CAGTTTGAC | AATTCCCCAT | TACATTATTA | |
| AGGGACTAAG | TTTGCCCAGG | AGCAGTGACC | CATAACAACC | AATCAGCAGG | TATGATTTCAC | 1200 |
| TCCCTGATTC | AAACGGGTCC | TCGTCACTGG | GTATTGTTGG | TTAGTCGTCC | ATACTAAATG | |
| TGGTCACCTG | TTTAAAAGCA | AAACATCTTAT | TGGTTGCTAT | GGGTTACTGC | TTCTGGGCAA | 1260 |
| ACCAGTGGAC | AAATTTCGT | TTGTAGAATA | ACCAACGATA | CCCAATGACG | AAGACCCGTT | |
| AATGTGTGCC | TCATAGGGGG | GTTAGTGTGT | TGTGTA | CTGTTACTGC | TTTATTTCAT | 1320 |
| TTACACACGG | AGTATCCCCC | CAATCACACA | ACACATGACT | TATTTAACAT | AAATAAAAGTA | |
| TGTTACAAAA | AAAAAAA | | | | | |
| ACAATGTTT | TTTTTTTT | | | | | |

Fig. 2. (Continuation page 2, SEQ ID NO:2).

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MSRTRKVDSL LLLAIPGLAL LLLPNAYCAS CEPVRIPMCK SMPWNMTKMP NHLHHSTQAN 60
AIALAIEQFEG LLTTECSQDL LFFLCAMYAP ICTIDFQHEP IKPCKSVcer ARAGCEPILI 120
KYRHTWPESL ACEELPVYDR GVCISPEAIV TVEQGTD SMP DFSMDSNNGN CGSGREHCKC 180
KPMKATQKTY LKNNNYNYVIR AKVKEVKVKC HDATAIVEVK EILKSSLVNI PKDTVTLYTN 240
SGCLCPQLVA NEEYIIMGYE DKERTRLLL V EGSLAEKWRD RLAKKVKRWD QKLRRPRKSK 300
DPVAPIPNKN SNSRQARS

Figure 3. Deduced amino acid sequence of Xenopus frazzled protein. SEQ ID NO:3.

Figure 4. Nucleotide sequence of the full-length frazzled cDNA derived from the Xenopus organizer. The sense strand of the DNA on top (5' to 3' direction) and the antisense strand on the bottom line (opposite direction). SEQ ID NO:4.

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| GAATTCCCTT TCACACAGGA CTCCCTGGCAG AGGTGAATGG TTAGCCCTAT GGATTTGGTT CTTAAGGGAA AGTGTGTCCT GAGGACCGTC TCCACTTACC AATCGGGATA CCTAAACCAA | 60 |
| TGTTGATTT GACACATGAT TGATTGCTT CAGATAGGAT TGAAGGACTT GGATTTTAT ACAACAAAAA CTGTGTACTA ACTAACGAAA GTCTATCCTA ACTTCCTGAA CCTAAAAATA | 120 |
| CTAATTCTGC ACTTTAAAT TATCTGAGTA ATTGTTCATT TTGTATTGGA TGGGACTAAA GATTAAGACG TGAAAATTA ATAGACTCAT TAACAAAGTAA AACATAACCT ACCCTGATT | 180 |
| GATAAACTTA ACTCCTTGCT TTTGACTTGC CCATAAACTA TAAGGTGGGG TGAGTTGTAG CTATTTGAAT TGAGGAACGA AAACTGAACG GGTATTTGAT ATTCCACCCCC ACTCAACATC | 240 |
| TTGCTTTAC ATGTGCCAG ATTTCCCTG TATTCCCTGT ATTCCCTCTA AAGTAAGCCT AACGAAAATG TACACGGTC TAAAAGGGAC ATAAGGGACA TAAGGGAGAT TTCATTGGA | 300 |
| ACACATACAG GTTGGGCAGA ATAACAATGT CTCGAACAAG GAAAGTGGAC TCATTACTGC TGTGTATGTC CAACCCGTCT TATTGTTACA GAGCTTGTTC CTTTCACCTG AGTAATGACG | 360 |
| TACTGGCCAT ACCTGGACTG GCGCTTCTCT TATTACCAA TGCTTACTGT GCTTCGTGTG ATGACCGGTA TGGACCTGAC CGCGAAGAGA ATAATGGTT ACGAATGACA CGAACACAC | 420 |
| AGCCTGTGCG GATCCCCATG TGCAAATCTA TGCCATGGAA CATGACCAAG ATGCCCAACC TCGGACACGC CTAGGGTAGAC ACGTTACCTT GTACTGGTTC TACGGGTTGG | 480 |
| ATCTCCACCA CAGCACTCAA GCCAATGCCA TCCTGGCAAT TGAACAGTTT GAAGGTTTGC TAGAGGTGGT GTCGTGAGTT CGGTTACGGT AGGACCGTTA ACTTGTCTAA CTTCCAAACG | 540 |
| TGACCACTGA ATGTAGCCAG GACCTTTGT TCTTTCTGTG TGCCATGTAT GCCCCCATT ACTGGTAGCT TACATCGTC CTGAAAACA AGAAAGACAC ACGGTACATA CGGGGGTAAA | 600 |
| GTACCATCGA TTTCCAGCAT GAACCAATTAA AGCCTTGCAA GTCCGTGTGC GAAAGGGCCA CATGGTAGCT AAAGGTGCTA CTTGGTTAAT TCGGAAOGTT CAGGCACACG CTTTCCCGGT | 660 |
| GGGCCGGCTG TGAGCCCATT CTCATAAAAGT ACCGGCACAC TTGGCCAGAG AGCCTGGCAT CCCGGCCGAC ACTCGGGTAA GAGTATTTCA TGGCCGTGTG AACCGGTCTC TCGGACCGTA | 720 |
| GTGAAGAGCT GCCCCGTATAT GACAGAGGAG TCTGCATCTC CCCAGAGGCT ATCGTCACAG CACTTCTCGA CGGGCATATA CTGTCCTCTC AGACGTAGAG GGGTCTCCGA TAGCAGTGTG | 780 |
| TGGAACAAGG AACAGATCA ATGCCAGACT TCTCCATGGA TTCAAACAAT GGAAATTGCG ACCTTGTCTCC TTGTCTAAGT TACGGTCTGA AGAGGTACCT AAGTTGTTA CCTTTAACGC | 840 |
| GAAGCGGCAG GGAGCACTGT AAATGCAAGC CCATGAAGGC AACCCAAAAG ACGTATCTCA CTTCGCCGTC CCTCGTGACA TTTACGTTCG GGTACTTCCG TTGGGTTTC TGCATAGAGT | 900 |
| AGAATAATTA CAATTATGTA ATCAGAGCAA AAGTGAAGA GGTGAAGTG AAATGCCACG TCTTATTAAT GTTAATACAT TAGTCTCGTT TTCACTTCTC CCACTTCAC TTTACGGTGC | 960 |
| ACGCAACAGC AATTGTGGAA GTAAAGGAGA TTCTCAAGTC TTCCCTAGTG AACATTCTA TGCCTGTGCG TTAACACCTT CATTCCCTCT AAGAGTTCAAG AAGGGATCAC TTGTAAGGAT | 1020 |

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| AAGACACAGT GACACTGTAC ACCAACTCAG GCTGCTTGTG CCCCCAGCTT GTGCCAATG TTCTGTGTCA CTGTGACATG TGGTTGAGTC CGACGAACAC GGGGGTCGAA CAACGGTTAC | 1080 |
| AGGAATACAT AATTATGGGC TATGAAAGACA AAGAGCGTAC CAGGCTTCTA CTAGTGGAAAG TCCTTATGTA TTAATACCCG ATACTTCTGT TTCTCGCATG GTCCGAAGAT GATCACCTTC | 1140 |
| GATCCTTGGC CGAAAAATGG AGAGATCGTC TTGCTAAGAA AGTCAAGCGC TGGGATCAAA CTAGGAACCG GCTTTTACCG TCTCTAGCAG AACGATTCTT TCAGTTCGCG ACCCTAGTTT | 1200 |
| AGCTTCGACG TCCCAGGAAA AGCAAAGACC CCGTGGCTCC AATTCCAAC AAAAACAGCA TCGAAGCTGC AGGGTCCTT TCGTTCTGG GGCACCGAGG TTAAGGGTTG TTTTGTGCGT | 1260 |
| ATTCCAGACA AGCGCGTAGT TAGACTAACG GAAAGGTGTA TGGAAACTCT ATGGACTTTG TAAGGTCTGT TCGCGCATCA ATCTGATTGC CTTTCCACAT ACCTTGAGA TACCTGAAAC | 1320 |
| AAACTAAGAT TTGCATTGTT GGAAGAGCAA AAAAGAAATT GCACTACAGC ACGTTATATT TTGATTCTA AACGTAACAA CCTTCTCGTT TTTTCTTTAA CGTGATGTCG TGCAATATAAA | 1380 |
| CTATTGTTA CTACAAGAAG CTGGTTAGT TGATTGTAGT TCTCCTTCC TTCTTTTTT GATAACAAAT GATGTTCTTC GACCAATCA ACTAACATCA AGAGGAAAGG AAGAAAAAAA | 1440 |
| TTATAACTAT ATTGCACGT GTTCCCAGGC AATTGTTTA TTCAACTTCC AGTGACAGAG AATATTGATA TAAACGTGCA CAAGGGTCCG TTAACAAAAT AAGTTGAAGG TCACTGTC | 1500 |
| CAGTGACTGA ATGTCTCAGC CAAAGAACG TCAATTCAATT TCTGATCAAC TAATGGTGAC GTCACTGACT TACAGAGTCG GATTCTTCG AGTTAAGTAA AGACTAGTTG ATTACCACTG | 1560 |
| AAGTGTGTTGA TACTTGGGGA AAGTGAACTA ATTGCAATGG TAAATCAGAG AAAAGTTGAC TTCACAAACT ATGAACCCCT TTCACTTGAT TAACGTTACC ATTTAGTCTC TTTCAACTG | 1620 |
| CAATGTTGCT TTTCTGTAG ATGAACAAGT GAGAGATCAC ATTTAAATGA TGATCACTTT GTTACAACGA AAAGGACATC TACTTGTCA CTCTCTAGTG TAAATTACT ACTAGTGAAA | 1680 |
| CCATTTAATA CTTTCAGCAG TTTTAGTTAG ATGACATGTA GGATGCACCT AAATCTAAAT GGTAAATTAT GAAAGTCGTC AAAATCAATC TACTGTACAT CCTACGTGGA TTTAGATTTA | 1740 |
| ATTTTATCAT AAATGAAGAG CTGGTTAGA CTGTATGGTC ACTGTTGGGA AGGTAAATGC TAAAATAGTA TTTACTTCTC GACCAATCT GACATACCAAG TGACAACCCCT TCCATTTACG | 1800 |
| CTACTTGTC AATTCTGTT TAAAAATTGC CAAATAAAT ATTAAGTCCT AAATAAAAAA GATGAAACAG TTAAGACAAA ATTTTTAACG GATTTATTTA TAATTCAAGGA TTTATTTTTT | 1860 |
| AAAAAAAAAA AAAAA TTTTTTTTT TTTT | |

Fig. 4. (Continuation page 2, SEQ ID NO:4).

MLLLFRAlPM LLLGLMVLQT DCEIAQYYID EEEPPGTVIA VLSQHSIFNT TDIPATNFRL 60
 MKQFNNSLIG VRESDGQLSI MERIDREQIC RQSLHCNLAL DVVSFSKGHF KLLNVKVEVR
 DINDHSPHFP SEIMHVEVSE SSSVGTRIPL EIAIDEDVGS NSIQNFQISN NSHFSIDVLT 120
 RADGVKYADL VLMRELDREI QPTYIMELLA MDGGVPSLSC TAVVNIRVLD FNDNSPVFER
 STIAVDLVED APLGYLLLEL HATDDDEGVN GEIVYGFSTL ASQEVRQLFK INSRTGSVTL 180
 EGQVDFETKQ TYEFEVQAQD LGPNPLTATC KVTVHILDVN DNTPAITITP LTTVNAGVAY 240
 IPETATKENF IALISTTDRA SGSGNGQVRCT LYGEHEHKLQ QAYEDSYMIV TTSTLDRENI 300
 AAYSLTVVAE DLGFPSLKTK KYYTVKVSDE NDNAPVFSKP QYEASILENN APGSYITTVI 360
 ARDSDDSDQNG KVNYRLVDAK VMGQSLTTFV SLDAADSGVLR AVRSLDYEKL KQLDFEIEAA 420
 DNGIPQLSTR VQLNLRIVDQ NDNCPVITNP LLNNNGSGEVL LPISAPQNYL VFQLKAEDSD 480
 EGHNSQLFYT ILRDPSRLFA INKESGEVFL KKQLNSDHSE DLSIVVAVYD LGRPSLSTNA 540
 TVKFILTDSE PSNVEVVILQ PSAEEQHQID MSIIFIAVLA GGCALLLLAI FFVACTCKKK 600
 AGEFKQVPEQ HGTCNEERLL STPSPQSVCSS SLSQSESQL SINTESENCS VSSNQEQQHQO 660
 TGIKHSISVP SYHTSGWHLD NCAMSISGHS HMGHISTKVQ WAKEIVTSMT VTLILVENQK 720
 RRALSSQCRH KPVLNTQMNQ QGSDMPITIS ATESTRVQKM GTAHCNMKRA IDCLTL 780
 840

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Figure 5. Deduced amino acid sequence of the Xenopus PAPC (paraxial protocadherin) protein. It encodes a member of the cadherin family of transmembrane proteins that has dorsalizing activity when constructs are injected into Xenopus embryos. SEQ ID NO:5.

Figure 6. Nucleotide sequence of the full-length PAPC cDNA derived from the *Xenopus* organizer. The sense strand of the DNA is shown in the top line (in the 5' to 3' direction), and the bottom line shows the antisense strand (opposite orientation). SEQ ID NO:6.

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| GAATTCCAG AGATGAACTC CTTGAGATTG TTTAAATGA CTGCAGGTCT GGAAGGGATT | 60 |
| CTTAAGGGTC TCTACTTGAG GAACTCTAAC AAAATTTACT GACGTCCAGA CCTTCCTAAC | |
| ACATTGCCAC ACTGTTCTA GGCAATGAAAA AACTGCAAGT TTCAACTTTG TTTTGGTGC | 120 |
| TGTAACGGTG TGACAAAGAT CCGTACTTTT TTGACGTTCA AAGTTGAAAC AAAAACCACG | |
| AACTTGATT CTTCAAGATG CTGCTCTCT TCAGAGCCAT TCCAATGCTG CTGTTGGAC | 180 |
| TTGAAACTAA GAAGTTCTAC GACGAAGAGA AGTCTCGGTAGGTTACGAC GACAACCCCTG | |
| TGATGGTTT ACAAACAGAC TGTGAAATTG CCCAGTACTA CATAGATGAA GAAGAACCCC | 240 |
| ACTACCAAAA TGTTGTCTG ACACCTAAC GGGTCATGAT GTATCTACTT CTTCTGGGG | |
| CTGGCACTGT AATTGCACTG TTGTCACAAC ACTCCATATT TAACACTACA GATATACCTG | 300 |
| GACCGTGACA TTAACGTCAAC AACAGTGTG TGAGGTATAA ATTGTGATGT CTATATGGAC | |
| CAACCAATTG CCGTCTAACG AAGCAATTAA ATAATTCCCT TATCGGAGTC CGTGAGAGTG | 360 |
| GTTGGTTAAA GGCAAGATTAC TTCGTTAAAT TATTAAGGGAA ATAGCCTCAG GCACTCTCAC | |
| ATGGGCAGCT GAGCATCATG GAGAGGATTG ACCGGGAGCA AATCTGCAGG CAGTCCCTTC | 420 |
| TACCCGTGCA CTCGTAGTAC CTCTCTAAC TGGCCCTCGT TTAGACGTCC GTCAGGGAAAG | |
| ACTGCAACCT GGCTTTGGAT GTGGTCAGCT TTCCCAAAGG ACACCTCAAG CTTCTGAACG | 480 |
| TGACGTTGGA CCGAAACCTA CACCAAGTCGA AAAGGTTCC TGTGAAGTTC GAAGACTTGC | |
| TGAAAGTGGG GGTGAGAGAC ATTAATGACC ATAGCCCTCA CTTTCCAGT GAAATAATGC | 540 |
| ACTTTCACCT CCACCTCTG TAATTACTGG TATCGGGAGT GAAAGGGTCA CTTTATTACG | |
| ATGTGGAGGT GTCTGAAAGT TCCTCTGTGG GCACCAAGGAT TCCTTAGAA ATTGCAATAG | 600 |
| TACACCTCCA CAGACTTCA AGGAGACACC CGTGGTCCTA AGGAAATCTT TAACTGTTATC | |
| ATGAAGATGT TGGGTCCAAC TCCATCCAGA ACTTTCAAGAT CTCAAATAAT AGCCACTTCA | 660 |
| TACTTCTACA ACCCAGGTTG AGGTAGGTCT TGAAAGTCTA GAGTTATTA TCGGTGAAGT | |
| GCATTGATGT GCTAACCAAGA GCAGATGGGG TGAAATATGC AGATTTAGTC TTAATGAGAG | 720 |
| CGTAACCTACA CGATTGGTCT CGTCTACCCCC ACTTTATACG TCTAAATCAG AATTACTCTC | |
| AACTGGACAG GGAAATCCAG CCAACATACA TAATGGAGCT ACTAGCAATG GATGGGGGTG | 780 |
| TTGACCTGTC CCTTTAGGTC GGTTGTATGT ATTACCTCGA TGATCGTTAC CTACCCCCAC | |
| TACCATCACT ATCTGGTACT GCAGTGGTTA ACATCCGAGT CCTGGACTTT AATGATAACA | 840 |
| ATGGTAGTGA TAGACCATGA CGTCACCAAT TGTAGGCTCA GGACCTGAAA TTACTATTGT | |
| GCCCCAGTGTGTT TGAGAGAAGC ACCATTGCTG TGGACCTAGT AGAGGATGCT CCTCTGGGAT | 900 |
| CGGGTCACAA ACTCTCTCG TGGTAACGAC ACCTGGATCA TCTCCTACGA GGAGACCCCTA | |
| ACCTTTGTT GGAGTTACAT GCTACTGACG ATGATGAAGG AGTGAATGGA GAAATTGTTT | 960 |
| TGAAAACAA CCTCAATGTA CGATGACTGC TACTACTTCC TCACTTACCT CTTAACAAA | |
| ATGGATTCAAG CACTTTGGCA TCTCAAGAGG TACGTCAAGCT ATTTAAAATT AACTCCAGAA | 1020 |
| TACCTAACGTC GTGAAACCGT AGAGTTCTCC ATGCAGTCGA TAAATTTAA TTGAGGTCTT | |

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| CTGGCAGTGT TACTCTTCAA GGCCAAAGTTG ATTTGAGAC CAAAGCAGACT TACGAATTG GACCGTCACA ATGAGAACCTT CCGGTTCAAC TAAAACCTCG GTTCGTCTGA ATGCTTAAAC | 1080 |
| AGGTACAAGC CCAAGATTTG GGCCCCAACC CACTGACTGC TACTTGAAA GTAAGTGTTC TCCATGTTCG GGTTCTAAC CGGGGGTTGG GTGACTGACG ATGAACATTT CATTGACAAG | 1140 |
| ATATACTTGA TGTAATGAT AATACCCCCAG CCATCACTAT TACCCCTCTG ACTACTGTAA TATATGAACT ACATTTACTA TTATGGGTC GGTAGTGATA ATGGGAGAC TGATGACATT | 1200 |
| ATGCAGGAGT TGCCTATATT CCAGAAACAG CCACAAAGGA GAACTTATA GCTCTGATCA TACGTCTCA ACGGATATAA GGTCTTGTC GGTGTTCCCT CTTGAAATAT CGAGACTAGT | 1260 |
| GCACTACTGA CAGAGCCTCT GGATCTAATG GACAAGTTCG CTGTACTCTT TATGGACATG CGTGATGACT GTCTCGGAGA CCTAGATTAC CTGTTCAAGC GACATGAGAA ATACCTGTAC | 1320 |
| AGCACTTAA ACTACAGCAA GCTTATGAGG ACAGTTACAT GATAGTTACC ACCTCTACTT TCGTGAAATT TGATGTCGTT CGAATACTCC TGTCAATGTA CTATCAATGG TGAGAGATGAA | 1380 |
| TAGACAGGGA AAACATAGCA GCGTACTCTT TGACAGTAGT TGCAGAAGAC CTTGGCTTCC ATCTGTCCTT TTTGTATCGT CGCATGAGAA ACTGTCATCA ACGTCTCTG GAACCGAAGG | 1440 |
| CCTCATTGAA GACCAAAAAG TACTACACAG TCAAGGTTAG TGATGAGAAAT GACAATGCAC GGAGTAACCTT CTGGTTTTC ATGATGTC AGTTCCAATC ACTACTCTTA CTGTTACGTG | 1500 |
| CTGTATTTTC TAAACCCAG TATGAAGCTT CTATTCTGGA AAATAATGCT CCAGGCTCTT GACATAAAAG ATTTGGGTC ATACTTCGAA GATAAGACCTT TTTATTACGA GGTCCGAGAA | 1560 |
| ATATAACTAC AGTGATAGCC AGAGACTCTG ATAGTGATCA AAATGGCAA GTAAATTACA TATATTGATG TCACTATCGG TCTCTGAGAC TATCACTAGT TTTACCGTTT CATTAAATGT | 1620 |
| GACTTGTGGA TGCAAAAGTG ATGGGCCAGT CACTAACAAAC ATTTGTTCTT CTTGATGCGG CTGAACACCT ACGTTTCAC TACCCGGTCA GTGATTGTTG TAAACAAAGA GAACTACGCC | 1680 |
| ACTCTGGAGT ATTGAGAGCT GTTAGGTCTT TAGACTATGA AAAACTTAAA CAACTGGATT TGAGACCTCA TAATCTCGA CAATCCAGAA ATCTGATACT TTTGAATT GTGACCTAA | 1740 |
| TTGAAATTGA AGCTGCAGAC AATGGGATCC CTCAACTCTC CACTCGCGTT CAACTAAATC AACTTTAATC TCGACGCTCG TTACCCCTAGG GAGTTGAGAG GTGAGCGCAA GTGATTAG | 1800 |
| TCAGAAATGT TGATCAAAAT GATAATTGCC CTGTGATAAC TAATCCTCTT CTTAATAATG AGTCTTATCA ACTAGTTTA CTATTAACGG GACACTATTG ATTAGGAGAA GAATTATTAC | 1860 |
| GCTCGGGTGA AGTTCTGCTT CCCATCAGCG CTCCTCAAAA CTATTTAGTT TTCCAGCTCA CGAGCCCAC TCAAGACGAA GGGTAGTCGC GAGGAGTTT GATAATCAA AAGGTCGAGT | 1920 |
| AAGCCGAGGA TTCAGATGAA GGGCACAACT CCCAGCTGTT CTATACCATA CTGAGAGATC TTCGGCTCCT AAGTCTACTT CCCGTGTTGA GGGTCGACAA GATATGGTAT GACTCTCTAG | 1980 |
| CAAGCAGATT GTTTGCCATT AACAAAGAAA GTGGTGAAAGT GTTCCGTAAA AAACAATTAA GTTCGTCTAA CAAACGGTAA TTGTTCTTT CACCACTTCA CAAGGACTTT TTTGTTAATT | 2040 |
| ACTCTGACCA TTCAGAGGAC TTGAGCATAG TAGTTGCACT GTATGACTTG GGAAGACCTT TGAGACTGGT AAGTCTCCTG AACTCGTATC ATCAACGTCA CATACTGAAC CCTTCTGGAA | 2100 |
| CATTATCCAC CAATGCTACA GTTAAATTCA TCCTCACCGA CTCTTTCCCT TCTAACGTTG GTAATAGGTG GTTACGATGT CAATTAAAGT AGGAGTGGCT GAGAAAAGGA AGATTGCAAC | 2160 |

Fig. 6. (Continuation page 2, SEQ ID NO:6).

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| AAGTCGTTAT TTTGCACCA TCTGCAGAAG AGCAGCACCA GATCGATATG TCCATTATAT TTCAGCAATA AAACGTTGGT AGACGTCTC TCGTCGTGGT CTAGCTATAC AGGTAATATA | 2220 |
| TCATTGCGAGT GCTGGCTGGT GGTTGTGCTT TGCTACTTTT GGCCATCTTT TTGTTGGCCT AGTAACGTCA CGACCGACCA CCAACACGAA ACGATGAAAA CCGGTAGAAA AAACACCGGA | 2280 |
| GTACTTGTAAGA AAAGAAAAGCT GGTGAATTAA ACCAGGTACC TGAACAACAC GGAACATGCA CATGAACATT TTTCTTCGA CCACCTAAAT TCGTCCATGG ACTTGTGTTG CCTTGTACGT | 2340 |
| ATGAAGAACG CCTGTTAACG ACCCCATCTC CCCAGTGGT CTCTCTCTC TTGTCAGT TACTTCTTGC GGACAATTG TGTTGAGAG GGGTCAGCCA GAGAAGAAGA AACAGAGTCA | 2400 |
| CTGAGTCATG CCAACTCTCC ATCAAATCTG AATCTGAGAA TTGCAGCGTG TCCTCTAAC GACTCAGTAC GGTTGAGAGG TAGTTATGAC TTAGACTCTT AACGTCGCAC AGGAGATTGG | 2460 |
| AAGAGCAGCA TCAGCAAACA GGCATAAAAGC ACTCCATCTC TGTACCATCT TATCACACAT TTCTCGTCGT AGTCGTTGT CCGTATTCG TGAGGTAGAG ACATGGTAGA ATAGTGTGTA | 2520 |
| CTGGTTGGCA CCTGGACAAT TGTGCAATGA GCATAAGTGG ACATTCTCAC ATGGGGCACA GACCAACCGT GGACCTGTTA ACACGTTACT CGTATTCAAC TGTAAGAGTG TACCCGTGT | 2580 |
| TTAGTACAAA GGTACAGTGG GCAAAGGAGA TAGTGACTTC AATGACAGTG ACTCTGATAC AATCATGTTT CCATGTCACC CGTTCTCTC ATCACTGAAG TTACTGTCAC TGAGACTATG | 2640 |
| TAGTGGAGAA TCAGAAAAGA AGAGCATTGA GCAGCCAATG CAGGCACAAG CCAGTGCTCA ATCACCTCTT AGTCTTCTC TCTCGTAAC CGTCGGTTAC GTCCGTGTTG GTCACGAGT | 2700 |
| ATACACAGAT GAATCAGCAG GGTTCCGACA TGCCGATAAC TATTCAGCC ACCGAATCAA TATGTGTCTA CTTAGTCGTC CCAAGGCTGT ACGGCTATTG ATAAAGTCGG TGGCTTAGTT | 2760 |
| CAAGGGTCCA GAAAATGGGA ACTGCACATT GCAATATGAA AAGGGCTATA GACTGTCTTA GTTCCCAGGT CTTTACCCCT TGACGTGTAA CGTTATACTT TTCCCGATAT CTGACAGAAT | 2820 |
| CTCTGTAGCT CCTGTATATT ACAATACCTA CCATGCAAGA ATGCCAACC TGCACATACC GAGACATCGA GGACATATAA TGTTATGGAT GGTACGTTCT TACGGATTGG ACGTGTATGG | 2880 |
| GAACCATACC CTTAGAGACC CTTATTACCA TATCAATAAT CCTGTTGCTA ATCGGATGCA CTTGGTATGG GAATCTCTGG GAATAATGGT ATAGTTATTA GGACAAAGAT TACCTACGT | 2940 |
| GGCGGAATAT GAAAGAGATT TAGTCAACAG AAGTGCAACG TTATCTCCGC AGAGATCGTC CCGCCTTATA CTTTCTCTAA ATCAGTTGTC TTCACGTTGC AATAGAGGGC TCTCTAGCAG | 3000 |
| TAGCAGATAC CAAGAATTCA ATTACAGTCC GCAGATATCA AGACAGCTTC ATCCTTCAGA ATCGTCTATG GTTCTTAAGT TAATGTCAGG CGTCTATAGT TCTGTGAAAG TAGGAAGTCT | 3060 |
| AATTGCTACA ACCTTTTAAT CATTAGGCAT GCAAGTGAGA ATGCACAAAG GCAAGTGCTT TTAACGATGT TGGAAAATTA GTAATCCGTA CGTTCACTCT TACGTGTTTC CGTTCACGAA | 3120 |
| TAGCATGAAA GCTAAATATA TGGAGTCTCC CCTTTCCCTC TGATGGATGG GGGGAGACAC ATCGTACTTT CGATTTATAT ACCTCAGAGG GGAAAGGGAG ACTACCTACC CCCCTCTGTG | 3180 |
| AGGACAGTGC ATAAATATAC AGCTGCTTTC TATTTGCATT TCACTGGGA ATTTTTGTT TCCTGTCAAG TATTTATATG TCGACGAAAG ATAAACGTAA AGTGAACCCCT TAAAAAACAA | 3240 |
| TTTTTACAT ATTTATTTT CCTGAATTGA ATGTGACATT GTCCTGTCAC CTAACTAGCA AAAAAAATGTA TAAATAAAAAA GGACTTAAC TACACTGTAA CAGGACAGTG GATTGATCGT | 3300 |

Fig. 6. (Continuation page 3, SEQ ID NO:6).

ATTAATCCA CAGACCTACA GTCAAATATT TGAGGGCCCC TGAAACAGCA CATCAGTCAG
TAATTTAGGT GTCTGGATGT CAGTTATAA ACTCCCGGGG ACTTTGTCGT GTAGTCAGTC 3360

GACCTAAAGT GGCTTTTA CTTTAGCAG CTCCCTGGTC TGCCCTCTGT GTTAATCAGC
CTGGATTCAGCGA AAAATCGTC GAGGACCCAG ACGGGAGACA CAATTAGTCG 3420

CCCTGGTCAA GTCTGAGTA GGATCATGGC GTTTTATAT GCATCTCACC TACTTGAC
GGGACCAAGTT CAGGACTCAT CCTAGTACCG CAAAAATATA CGTAGAGTGG ATGAAACCTG 3480

GTGATTTACA CATAATAGGA AACGCTTGGT TTCAGTGAAG TCTGTGTGT ATATATTCTG
CACTAAATGT GTATTATCCT TTGCGAACCA AAGTCACTTC AGACACAAACA TATATAAGAC 3540

TTATATACAC GCATTTGTG TTTGTGTATA TATTCAAGT CCATTCAAGAT ATGTGTATAT
AATATATGTG CGTAAAACAC AAACACATAT ATAAAGTTCA GGTAAGTCTA TACACATATA 3600

AGTGCAGACC TTGTAAATTA AATATTCTGA TACTTTTCC TCAATAATA TTTAAAT
TCACGTCTGG AACATTTAAT TTATAAGACT ATGAAAAGG AGTTATTAT AAATTTA

Fig. 6. (Continuation page 4, SEQ ID NO:6).

0660 0660 0660 0660 0660 0660 0660 0660

MVCCGPGRML LGWAGLLVLA ALCLLQVPGA QAAACEPVRI PLCKSLPWNN TKMPNHLHHS 60
TQANAILAME QFEGLLGTHC SPDLLFFLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE 120
PILIKYRHSHW PESLACDELP VYDRGVCISP EAIVTADGAD FPMDSSSTGHC RGASSERCKC 180
KPVRATQKTY FRNNNYNYVIR AKVKEVKMKC HDVTAVVEVK EILKASLVNI PRDTVNLYTT 240
SGCLCPPLTV NEEYVIMGYE DEERSRLLL EGSIAEKWKD RLGKKVKRWD MKLRHLGLGK 300
TDASDSTQNZ KSGRNSNPRP ARS.

Figure 7. Deduced amino acid sequence of mouse FRZB-1 protein. SEQ ID NO:7.

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Figure 8. Nucleotide sequence of the full-length mouse FRZB-1 cDNA. SEQ ID NO:8.

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|---|-----|
| AAGCCTGGGA CCATGGTCTG CTGCGGCCCG GGACGGATGC TGCTAGGATG GGCCGGGTIG TTCGGACCTT GGTACAGAC GACGCCGGC CCTGCCTACG ACGATCCTAC CGGGCCCAAC | 60 |
| CTAGTCTGG CTGCTCTCTG CCTGCTCCAG GTGCCCGGAG CTCAGGCTGC AGCCTGTGAG GATCAGGACC GACGAGAGAC GGACGAGGTC CACGGGCCTC GAGTCCGACG TCGGACACTC | 120 |
| CCTGTCCGCA TCCCCTGCTG CAAGTCCCTT CCCTGGAACA TGACCAAGAT GCCCAACCAC GGACAGGCGT AGGGCGACAC GTTCAGGGAA GGGACCTTGT ACTGGTTCTA CGGGTTGGTG | 180 |
| CTGCACCCACA GCACCCAGGC TAACGCCATC CTGGCCATGG AACAGTTCGA AGGGCTGCTG GACGTGGTGT CGTGGGTCCG ATTGCGGTAG GACCGGTACC TTGTCAAGCT TCCCAGCAC | 240 |
| GGCACCCACT GCAGCCCGGA TCTTCTCTTC TTCCTCTGTG CAATGTACGC ACCCATTTGC CCGTGGGTGA CGTCGGGCCT AGAAGAGAAC AGGAGACAC GTTACATGCG TGGTAAACG | 300 |
| ACCATCGACT TCCAGCACGA GCCCATCAAG CCCTGCAAGT CTGTGTGTGA GCGCGCCCGA TGGTAGCTGA AGGTCGTGCT CGGGTAGTTC GGGACGTTCA GACACACACT CGCGCGGGCT | 360 |
| CAGGGCTGCG AGCCCATTCT CATCAAGTAC CGCCACTCGT GGCCGGAAAG CTTGGCTGCG GTCCCGACGC TCAGGTAAGA GTAGTTCATG GCGGTGAGCA CCGGCCTTTC GAACCGGACG | 420 |
| GACGGAGCTGC CGGTGTACGA CCGCGGCGTG TGCATCTCTC CTGAGGCCAT CGTCACCGCG CTGCTCGACG GCCACATGCT GGCGCCGCAC ACGTAGAGAG GACTCCGGTA GCAGTGGCGC | 480 |
| GACGGAGCGG ATTTCTAT GGATTCAAGT ACTGGACACT GCAGAGGGGC AAGCAGCGAA CTGCCTCGCC TAAAAGGATA CCTAAGTTCA TGACCTGTGA CGTCTCCCCG TTCGTCGCTT | 540 |
| CGTGCCTAAAT GTAAGCCTGT CAGAGCTACA CAGAAGACCT ATTTCCGGAA CAATTACAAC GCAACGTTTA CATTGGACA GTCTCGATGT GTCTCTGGAA TAAAGGCTT GTTAATGTTG | 600 |
| TATGTCATCC GGGCTAAAGT TAAAGAGGTA AAGATGAAAT GTCATGATGT GACCGCCGTT ATACAGTAGG CCCGATTCA ATTTCTCCAT TTCTACTTTA CAGTACTACA CTGGCGGCAA | 660 |
| GTGGAAGTGA AGGAAATTCT AAAGGCATCA CTGGTAAACA TTCCAAGGGAA CACCGTCAAT CACCTTCACT TCCTTTAAGA TTCCGTAGT GACCATTGT AAGGTTCCCT GTGGCAGTTA | 720 |
| CTTTATACCA CCTCTGGCTG CCTCTGTCTT CCACTTACTG TCAATGAGGA ATATGTCATC GAAATATGGT GGAGACCGAC GGAGACAGGA GGTGAATGAC AGTTACTCCT TATACAGTAG | 780 |
| ATGGGCTATG AAGACGAGGA ACGTTCCAGG TTACTCTGG TAGAAGGCTC TATACTGAG TACCCGATAC TTCTGCTCCT TGCAAGGTCC AATGAGAACC ATCTTCCGAG ATATCGACTC | 840 |
| AAAGTGGAAAGG ATCGGCTTGG TAAGAAAGTC AAGCGCTGGG ATATGAAACT CCGACACCTT TTCACCTTCC TAGCCGAACC ATTCTTCAAG TTGCGACCC TATACTTTGA GGCTGTGGAA | 900 |
| GGACTGGGTA AAACTGATGC TAGCGATTCC ACTCAGAATC AGAAGTCTGG CAGGAACCTC CCTGACCCAT TTTGACTACG ATCGCTAAGG TGAGTCTTAG TCTTCAGACC GTCTTGAGA | 960 |

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| | |
|---|------|
| AATCCCCGGC CAGCACGCAG CTAATCCTG AAATGTAAAA GGCCACACCC ACGGACTCCC TTAGGGCGG GTCGTGCGTC GATTAGGAC TTTACATTT CCGGTGTGGG TGCGTGAGGG | 1020 |
| TTCTAAGACT GGCGCTGGTG GACTAACAAA GGAAAACCGC ACAGTTGTGC TCGTGACCGA AAGATTCTGA CCGCGACCAC CTGATTGTT CCTTTGGCG TGTCAACACG AGCACTGGCT | 1080 |
| TTGTTTACCG CAGACACCGC GTGGCTACCG AAGTTACTTC CGGTCCCCCTT TCTCCTGCTT AACAAATGGC GTCTGTGGCG CACCGATGGC TTCAATGAAG GCCAGGGGAA AGAGGACGAA | 1140 |
| CTTAATGGCG TGGGGTTAGA TCCTTTAATA TGTTATATAT TCTGTTTCAT CAATCACGTG GAATTACCGC ACCCCAATCT AGGAAATTAT ACAATATATA AGACAAAGTA GTTAGTGCAC | 1200 |
| GGGACTGTTC TTTTGCAACC AGAATAGTAA ATTAATATG TTGATGCTAA GGTTCTGTA CCCTGACAAG AAAACGTTGG TCTTATCATT TAATTATAC AACTACGATT CCAAAGACAT | 1260 |
| CTGGACTCCC TGGGTTTAAT TTGGTGTTC GTACCCGTAT TGAGAATGCA ATGTTTCATG GACCTGAGGG ACCCAAATTA AACCCACAAGA CATGGGACTA ACTCTTACGT TACAAAGTAC | 1320 |
| TAAAGAGAGA ATCCTGGTCA TATCTCAAGA ACTAGATATT GCTGTAAGAC AGCCTCTGCT ATTCTCTCT TAGGACCAAGT ATAGAGTTCT TGATCTATAA CGACATTCTG TCGGAGACGA | 1380 |
| GCTGCGCTTA TAGTCTTGTG TTTGTATGCC TTTGTCCATT TCCCTCATGC TGTGAAAGTT CGACGCGAAT ATCAGAACAC AAACATACGG AAACAGGTAA AGGGAGTACG ACACTTCAA | 1440 |
| ATACATGTTT ATAAAGGTAG AACGGCATTG TGAAATCAGA CACTGCACAA GCAGAGTAGC TATGTACAAA TATTTCATC TTGCCGTAAA ACTTTAGTCT GTGACGTGTT CGTCTCATCG | 1500 |
| CCAACACCAAG GAAGCATTG TGAGGAAACG CCACACAGCA TGACTTATTT TCAAGATTGG GGTTGTGGTC CTTCGTAAAT ACTCCTTTGC GGTGTGTCGT ACTGAATAAA AGTTCTAAC | 1560 |
| CAGGCAGCAA AATAAATAGT GTTGGGAGCC AAGAAAAGAA TATTTGCCT GGTTAAGGGG GTCCGTCGTT TTATTTATCA CAACCCTCGG TTCTTTCTT ATAAAACGGA CCAATTCCCC | 1620 |
| CACACTGGAA TCAGTAGCCC TTGAGCCATT AACAGCAGTG TTCTCTGGC AAGTTTTGA GTGTGACCTT AGTCATCGGG AACTCGGTAA TTGTCGTCAC AAGAAGACCG TTCAAAAAC | 1680 |
| TTTGTTCATA AATGTATTCA CGAGCATTAG AGATGAACCT ATAACTAGAC ATCTGTTGTT AAACAAGTAT TTACATAAGT GCTCGTAATC TCTACTTGAA TATTGATCTG TAGACAACAA | 1740 |
| ATCTCTATAG CTCTGCTTCC TTCTAAATCA AACCCATTGT TGGATGCTCC CTCTCCATTC TAGAGATATC GAGACGAAGG AAGATTTAGT TTGGGTAAACA ACCTACGAGG GAGAGGTAAG | 1800 |

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ATAAATAAAT TTGGCTGCT GTATTGGCCA GGAAAAGAAA GTATTAAAGT ATGCATGCAT 1860
TATTATTTA AACCGAACGA CATAACCGGT CCTTTCTTT CATAATTCA TACGTACGTA
GTGCACCAGG GTGTTATTTA ACAGAGGTAT GTAACTCTAT AAAAGACTAT AATTCACAGG 1920
CACGTGGTCC CACAATAAAT TGCTCCATA CATTGAGATA TTTCTGATA TTAAATGTCC
ACACGGAAAT GTGCACATTT GTTTACTTTT TTTCTCCTT TTGCTTGAG CTTGTGATTT 1980
TGTGCCTTTA CACGTGTAAA CAAATGAAAA AAAGAAGGAA AACGAAACCC GAACACTAAA
TGGTTTTGG TGTGTTATG TCTGTATTTT GGGGGTGGG TAGGTTTAAG CCATTGCACA 2040
ACCAAAAC ACCACAAATAC AGACATAAAA CCCCCCACCC ATCCAAATTC GGTAACGTGT
TTCAAGTTGA ACTAGATTAG AGTAGACTAG GCTCATGGC CTAGACATTA TGATTGAAT 2100
AAGTTCAACT TGATCTAATC TCATCTGATC CGAGTAACCG GATCTGTAAT ACTAAACTTA
TTGTGTTGT TAATGCTCCA TCAAGATGTC TAATAAAAGG AATATGGTG TCAACAGAGA 2160
AACACAAACAA ATTACGAGGT AGTTCTACAG ATTATTTCC TTATACCAAC AGTTGTCTCT
CGACAAACAAC AACAAA
GCTGTTGTTG TTGTTT

MVCGSPGGML LLRAGLLALA ALCLLRVPGA RAAACEPVRI PLCKSLPWNM TKMPNHLHHS 60
TQANAILAIE QFEGLLGTHC SPDLLFFLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE 120
PILIKYRHSH PENLACEELP VYDRGVCISP EAIVTADGAD FPMDSNGNC RGASSERCKC 180
KPIRATQKTY FRNNNYNYVIR AKVKEIKTKC HDVTAVVEVK EILKSSLVNI PRDTVNLYTS 240
SGCLCPPLNV NEEYIIMGYE DEERSRLLL V EGSI AEKW KD RLGKKV KRWD MKLRHLGLSK 300
SDSSNSDSTQ SQKSGRNSNP RQARN.

Figure 9. Dduced amino acid sequence of human FRZB-1 protein. SEQ ID NO:9.

0900000000 - 074104

Figure 10. Nucleotide sequence of the full-length human FRZB-1 cDNA. SEQ ID NO:10.
 This sequence was assembled from public ESTs from the Genbank database
 (accession numbers: H18848, R63748, W38677, W44760, H38379 and N71244).

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|---|-----|
| GGCGGAGCGG GCCTTTGGC GTCCACTGCG CGGCTGCACC CTGCCCATC TGCCGGGATC CGCCTCGCC CGGAAAACCG CAGGTGACGC GCCGACGTGG GACGGGGTAG ACGGCCCTAG | 60 |
| ATGGTCTGCG GCAGCCCGGG AGGGATGCTG CTGCTGCGGG CGGGCTGCT TGCCCTGGCT TACCAAGACGC CGTCGGGCC TCCCTACGAC GACGACGCC GGCCGACGA ACGGGACCGA | 120 |
| GCTCTCTGCC TGCTCCGGGT GCCCGGGGCT CGGGCTGCAG CCTGTGAGCC CGTCCGCATC CGAGAGACGG ACGAGGCCA CGGGCCCCGA GCCCGACGTC GGACACTCGG GCAGGCGTAG | 180 |
| CCCTGTGCA AGTCCCTGCC CTGGAACATG ACTAAAGATGC CCAACCACCT GCACCACAGC GGGGACACGT TCAGGGACGG GACCTTGAC TGATTCTACG GGTTGGTGGA CGTGGTGTG | 240 |
| ACTCAGGCCA ACGCCATCCT GGCCATCGAG CAGTCGAAG GTCTGCTGGG CACCCACTGC TGAGTCCGGT TGCGGTAGGA CCGGTAGCTC GTCAAGCTTC CAGACGACCC GTGGGTGACG | 300 |
| AGCCCCGATC TGCTCTTCTT CCTCTGTGCC ATGTACGCGC CCATCTGCAC CATTGACTTC TCGGGGCTAG ACGAGAAAGAA GGAGACACGG TACATGCGCG GGTAGACGTG GTAAGTGAAG | 360 |
| CAGCACGAGC CCATCAAGCC CTGTAAGTCT GTGTGCGAGC GGGCCCGCA GGGCTGTGAG GTCGTGCTCG GGTAGTTCGG GACATTAGA CACACGCTCG CCCGGCCGT CCCGACACTC | 420 |
| CCCATACTCA TCAAGTACCG CCACTCGTGG CCGGAGAACCC TGGCCTGCGA GGAGCTGCCA GGGTATGAGT AGTTCATGGC GGTGAGCACC GGCCTCTTGG ACCGGACGCT CCTCGACGGT | 480 |
| GTGTACGACA GGGCGTGTG CATCTCTCCC GAGGCCATCG TTACTGCGGA CGGAGCTGAT CACATGCTGT CCCCCCACAC GTAGAGAGGG CTCCGGTAGC AATGACGCC GCCTCGACTA | 540 |
| TTTCCTATGG ATTCTAGTAA CGGAAACTGT AGAGGGCAA GCAGTGAACG CTGTAAATGT AAAGGATACC TAAGATCATT GCCTTTGACA TCTCCCCGTT CGTCACTTGC GACATTACCA | 600 |
| AAGCCTATTA GAGCTACACA GAAGACCTAT TTCCGGAACA ATTACAACCA TGTCATTGG TTCGGATAAT CTCGATGTGT CTTCTGGATA AAGGCCTTGT TAATGTTGAT ACAGTAAGCC | 660 |
| GCTAAAGTTA AAGAGATAAA GACTAAGTGC CATGATGTGA CTGCAGTAGT GGAGGTGAAG CGATTCAAT TTCTCTATTT CTGATTACG GTACTACACT GACGTCACTCA CCTCCACTTC | 720 |
| GAGATTCTAA AGTCCTCTCT GGTAAACATT CCACGGGACA CTGTCAACCT CTATACCAGC CTCTAAGATT TCAGGAGAGA CCATTGTAA GGTGCCCTGT GACAGTTGGA GATATGGTGG | 780 |
| TCTGGCTGCC TCTGCCCTCC ACTTAATGTT AATGAGGAAT ATATCATCAT GGGCTATGAA AGACCGACGG AGACGGGAGG TGAATTACAA TTACTCCTTA TATAGTAGTA CCCGATACTT | 840 |

GATGAGGAAC GTTCCAGATT ACTCTGGTG GAAGGCCTCA TAGCTGAGAA GTGGAAGGAT 900
 CTACTCCTTG CAAGGTCTAA TGAGAACAC CTTCCGAGAT ATCGACTCTT CACCTTCCTA

 CGACTCGGTA AAAAGTTAA GCGCTGGGAT ATGAAGCTTC GTCATCTTGG ACTCAGTAAA 960
 GCTGAGCCAT TTTTCATT CGCGACCCTA TACTTCGAAG CAGTAGAAC TGAGTCATT

 AGTGATTCTA GCAATAGTGA TTCCACTCAG AGTCAGAAGT CTGGCAGGAA CTCGAACCCC 1020
 TCACTAAGAT CGTTATCACT AAGGTGAGTC TCAGTCTTCA GACCGTCCCTT GAGCTTGGGG

 CGGCAAGCAC GCAACTAAAT CCCGAAATAC AAAAGTAAC ACAGTGGACT TCCTATTAAG 1080
 GCCGTTCGTG CGTTGATTAA GGGCTTTATG TTTTCATTG TGTCACCTGA AGGATAATTG

 ACTTACTTGC ATTGCTGGAC TAGCAAAGGA AAATTGCACT ATTGCACATC ATATTCTATT 1140
 TGAATGAACG TAACGACCTG ATCGTTCCCT TTTAACGTGA TAACGTGTAG TATAAGATAA

 GTTTACTATA AAAATCATGT GATAACTGAT TATTACTTCT GTTTCTCTTT TGGTTTCTGC 1200
 CAAATGATAT TTTTAGTACA CTATTGACTA ATAATGAAGA CAAAGAGAAA ACCAAAGACG

 TTCTCTCTTC TCTCAACCCC TTTGTAATGG TTTGGGGCA GACTCTTAAG TATATTGTGA 1260
 AAGAGAGAAG AGAGTTGGGG AAACATTACC AAACCCCCGT CTGAGAATTG ATATAACACT

 GTTTCTATT TCACTAATCA TGAGAAAAAC TGTTCTTTG CAATAATAAT AAATTAAACA 1320
 CAAAGATAA AGTGATTAGT ACTCTTTTG ACAAGAAAAC GTTATTATTA TTTAATTGT

 TGCTGTTACC AGAGCCTCTT TGCTGAGTCT CCAGATGTTA ATTTACTTTG TGCAACCCAA 1380
 ACGACAATGG TCTCGGAGAA ACGACTCAGA GGTCTACAAT TAAATGAAAG ACGTGGGGTT

 TTGGGAATGC AATATTGGAT GAAAAGAGAG GTTTCTGGTA TTCACAGAAA GCTAGATATG 1440
 AACCCCTACG TTATAACCTA CTTTTCTCTC CAAAGACCCT AAGTGTCTTT CGATCTATAC

 CCTTAAAACA TACTCTGCCG ATCTAATTAC AGCCTTATTG TTGTATGCCT TTTGGGCATT 1500
 GGAATTTGT ATGAGACGGC TAGATTAAATG TCGGAATAAA AACATACGGA AAACCCGTAA

 CTCCTCATGC TTAGAAAGTT CCAAATGTTT ATAAAGGTAA AATGGCAGTT TGAAGTCAAA 1560
 GAGGAGTACG AATCTTCAA GGTTTACAAA TATTCACATT TTACCGTCAA ACTTCAGTTT

 TGTCACATAG GCAAAGCAAT CAAGCACCAG GAAGTGTGTTA TGAGGAAACA ACACCCAAGA 1620
 ACAGTGTATC CGTTCTGTTA GTTCGTGGTC CTTCACAAAT ACTCCTTGT TGTGGTTCT

 TGAATTATTT TTGAGACTGT CAGGAAGTAA AATAAATAGG AGCTTAAGAA AGAACATTTT 1680
 ACTTAATAAA AACTCTGACA GTCCTTCATT TTATTTATCC TCGAATTCTT TCTTGTAAAA

 GCCTGATTGA GAAGCACAAC TGAAACCAGT AGCCGCTGGG GTGTTAATGG TAGCATTCTT 1740
 CGGACTAACT CTTCGTGTG ACTTTGGTCA TCGGCGACCC CACAATTACC ATCGTAAGAA

 CTTTTGGCAA TACATTGAT TTGTTCATGA ATATATTAAT CAGCATTAGA GAAATGAATT 1800
 GAAAACCGTT ATGTAACACTA AACAAAGTACT TATATAATTA GTCGTAATCT CTTTACTTAA

 ATAACCTAGAC ATCTGCTGTT ATCACCATAG TTTGTTTAA TTTGCTTCCT TTTAAATAAA 1860
 TATTGATCTG TAGACGACAA TAGTGGTATC AAAACAAATT AAACGAAGGA AAATTTATTT

 CCCATTGGTG AAAGTCAAAA AAAAAAAA AAA
 GGGTAACCAC TTTCAGTTT TTTTTTTTT TTT

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